

Table 7. Primers

SEQ ID NO. 96	GPT	5'-GTTTAACCACTGCATGATG-3'
SEQ ID NO. 97	TK _R	5'-GTCCTCATCGAGTGGCTAC-3'
SEQ ID NO. 98	GLY1	5'-CGTGACATGGTACATTCGGACACTGGGCACATTAAGCGGA-3'
5	SEQ ID NO. 99	5'-TGCCTCATACACAATGGAGCTCTGGGACGAGTCGTTCTGTGAC-3'
SEQ ID NO. 100	GLY2	5'-TACCCAGCAGGGG <u>GAGC</u> TCGTTGCTCCGAAACGCAGGGCAC-3'
SEQ ID NO. 101	GLY3	5'-TGTCTGGTGGGGACGG <u>AGGC</u> CTAGCTGCAGGCTGGG-3'
SEQ ID NO. 102	GLY4	5'-CGTTATGTGG <u>CCGGGT</u> AGATTGAGGACTGGCAGTCCTGCACCGTCTC-3'
SEQ ID NO. 103	GLY5	5'-CAGGGCCGTTG <u>AGGC</u> CTCCACTGCATCATATCCAAAGC-3'
SEQ ID NO. 104	OVR1	5'-CGGG <u>GA</u> ATGTACCATGTCACGAACGAC-3'
SEQ ID NO. 105	OVR2	5'- <u>GCTC</u> CATTG <u>GT</u> TATGAGGCAGGG-3'
SEQ ID NO. 106	OVR3	5'- <u>GAGC</u> T <u>CCG</u> GCTGGTAGGCC-3'
SEQ ID NO. 107	OVR4	5'- <u>CC</u> TC <u>GG</u> CC <u>CC</u> AC <u>CC</u> AC <u>GA</u> AA <u>TA</u> ACG-3'
SEQ ID NO. 108	OVR5	5'-CTAC <u>CC</u> GG <u>CC</u> AC <u>AT</u> ACGG <u>GT</u> CA <u>CC</u> G-3'
SEQ ID NO. 109	OVR6	5'-GGAG <u>GG</u> CTAC <u>AA</u> ACGG <u>CC</u> CTGG <u>GT</u> G-3'
SEQ ID NO. 110	GPT-2	5'-TTCTAT <u>CG</u> ATTA <u>AA</u> ATGA <u>ATTC</u> -3'
SEQ ID NO. 111	TK _R -2	5'-GCCATACG <u>CT</u> ACAG <u>CC</u> GAT <u>CC</u> -3'

nucleotides underlined represent additional restriction site
nucleotides in bold represent mutations with respect to the original HCC110A sequence

original paper
copy,
highlighted bold
letters.

Table 7. Primers

SEQ ID NO. 96	GPT	5'- GTTTAA CCACTGCATGATG-3'
SEQ ID NO. 97	TK _n	5'- GTC CCATCGAGTGC GGCTAC-3'
SEQ ID NO. 98	GLY1	5'- CGT GACATGGTACAT CCGG ACACTTGGCGACATTAAGCGGA-3'
SEQ ID NO. 99	GLY2	5'- TG CCCTACACAA ATGGAGCT GGGACGAGTGGTGGAC-3'
SEQ ID NO. 100	GLY3	5'- TACCC CAGGGGAG GCTCT GTGGCTCCCGAACCGAGGGCAC-3'
SEQ ID NO. 101	GLY4	5'- TGTGG TGGGGAGGG GGCT GCCTAGTGGGAGGGTGGG-3'
SEQ ID NO. 102	GLY5	5'- CGTT ATGTGGCC GGGG TAGATTGAGC ACTGG CAGTCTGCACCGGTCTC-3'
SEQ ID NO. 103	GLY6	5'- CAGGGCCGTTGTA GG CCACTGCATCATCATATCCC AAGC-3'
SEQ ID NO. 104	OVR1	5'- CCGGAA TGTACCCATGTCA CGAACGAC -3'
SEQ ID NO. 105	OVR2	5'- GCTCC ATTGTGTATGAGGG ACGGG -3'
SEQ ID NO. 106	OVR3	5'- GAGG <u>CC</u> GGCTGGGTAGGG GGG -3'
SEQ ID NO. 107	OVR4	5'- GGTCCC GTCC CCAC ACGACAAATACG-3'
SEQ ID NO. 108	OVR5	5'- CTACCCGGG GCACATAACGGGG TCACCG -3'
SEQ ID NO. 109	OVR6	5'- GGAGGG CTACAA ACGGCC CTGGTGG-3'
SEQ ID NO. 110	GPT-2	5'- TTCT TATCGATTAA ATGA ATTTC-3'
SEQ ID NO. 111	TK _n -2	5'- GCCATACG GCTCACAGCCGATCCC-3'

nucleotides in bold represent additional mutations with respect to the original HICCH10A sequence
nucleotides in bold represent additional restriction site

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